

GenCore version 4.5
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OM protein - protein search, using sw model

Run on: December 26, 2001, 10:34:59 ; Search time 98.13 Seconds
(without alignments)
1324.197 Million cell updates/sec

Title: US-09-497-967-7
Perfect score: 2540
Sequence: 1 MKNILVILISLFINQIKS.....QCDNFANFLISLLLSIYYLL 468

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 3148936 seqs, 277657034 residues

Total number of hits satisfying chosen parameters: 3148936

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Pending_Patents_AA_Main:*

1: /cgn2_6/ptodata/2/paa/PCTUS_COMB.pep.*
2: /cgn2_6/ptodata/2/paa/US06_COMB.pep.*
3: /cgn2_6/ptodata/2/paa/US07_COMB.pep.*
4: /cgn2_6/ptodata/2/paa/US080_COMB.pep.*
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6: /cgn2_6/ptodata/2/paa/US082_COMB.pep.*
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23: /cgn2_6/ptodata/2/paa/US099_COMB.pep.*
24: /cgn2_6/ptodata/2/paa/US60_COMB.pep.*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	2540	100.0	468	18	US-09-498-612-6
2	921	36.3	442	18	US-09-498-612-5
3	843.5	33.2	414	3	US-07-763-352A-15
4	749	29.5	375	3	US-07-763-352A-3
5	251	9.9	3131	1	PCT-US01-06960-2
6	251	9.9	3131	19	US-09-515-363-2
7	251	9.9	3131	19	US-09-515-363B-2
8	245.5	9.7	3880	4	US-08-028-021-1
9	242.5	9.5	544	21	US-09-791-932-40
					Sequence 6, Appli
					Sequence 5, Appli
					Sequence 15, Appli
					Sequence 3, Appli
					Sequence 2, Appli
					Sequence 2, Appli
					Sequence 1, Appli
					Sequence 40, Appli

10	238.5	9.4	2826	1	PCT-US97-17746-50	Sequence 50, Appli
11	238.5	9.4	2826	16	US-09-254-776-50	Sequence 50, Appli
12	237.5	9.4	914	1	PCT-US00-08561-47	Sequence 47, Appli
13	233	9.2	1917	18	US-09-436-063-5	Sequence 5, Appli
14	233	9.2	1917	18	US-09-436-063C-5	Sequence 5, Appli
15	233	9.2	1917	20	US-09-627-650-5	Sequence 5, Appli
16	233	9.2	1917	20	US-09-627-650A-5	Sequence 5, Appli
17	233	9.2	1917	20	US-09-627-650B-5	Sequence 5, Appli
18	230.5	9.1	480	22	US-09-823-936-16	Sequence 16, Appli
19	229.5	9.0	1277	1	PCT-US99-19726-33	Sequence 33, Appli
20	229.5	9.0	1277	19	US-09-512-255-33	Sequence 33, Appli
21	228.5	9.0	426	18	US-09-413-198-500	Sequence 500, Appli
22	228	9.0	474	6	US-08-260-670-15	Sequence 15, Appli
23	224	8.8	575	21	US-09-791-932-46	Sequence 46, Appli
24	222	8.7	765	17	US-09-308-823A-484	Sequence 484, Appli
25	221	8.7	105	15	US-09-196-161D-1	Sequence 1, Appli
26	221	8.7	105	15	US-09-196-161D-10	Sequence 10, Appli
27	220	8.7	505	6	US-08-260-670-16	Sequence 16, Appli
28	219	8.6	3334	18	US-09-478-081-757	Sequence 757, Appli
29	217.5	8.6	2443	18	US-09-478-081-1011	Sequence 1011, Appli
30	217	8.5	2508	18	US-09-436-063-7	Sequence 7, Appli
31	217	8.5	2508	18	US-09-436-063C-7	Sequence 7, Appli
32	217	8.5	2508	20	US-09-627-650-7	Sequence 7, Appli
33	217	8.5	2508	20	US-09-627-650A-7	Sequence 7, Appli
34	217	8.5	2508	20	US-09-627-650B-7	Sequence 7, Appli
35	217	8.5	2544	18	US-09-436-063-3	Sequence 3, Appli
36	217	8.5	2544	18	US-09-436-063C-3	Sequence 3, Appli
37	217	8.5	2544	20	US-09-627-650-3	Sequence 3, Appli
38	217	8.5	2544	20	US-09-627-650A-3	Sequence 3, Appli
39	217	8.5	2544	20	US-09-627-650B-3	Sequence 3, Appli
40	217	8.5	2601	18	US-09-436-063-9	Sequence 9, Appli
41	217	8.5	2601	18	US-09-436-063C-9	Sequence 9, Appli
42	217	8.5	2601	20	US-09-627-650-9	Sequence 9, Appli
43	217	8.5	2601	20	US-09-627-650A-9	Sequence 9, Appli
44	217	8.5	2601	20	US-09-627-650B-9	Sequence 9, Appli
45	217	8.5	3518	19	US-09-522-097B-2	Sequence 2, Appli

ALIGNMENTS

RESULT 1
US-09-498-612-6
; Sequence 6, Application US/09498612
; GENERAL INFORMATION:
; APPLICANT: GAERTIG, Jacek
; APPLICANT: DICKERSON Jr., Harry W.
; APPLICANT: CLARK, Theodore G.
; APPLICANT: THE UNIVERSITY OF GEORGIA RESEARCH FOUNDATION, INC
; TITLE OF INVENTION: RECOMBINANT EXPRESSION OF HETEROLOGOUS NUCLEIC ACIDS IN
; TITLE OF INVENTION: PROTOZOA
; FILE REFERENCE: 235.00100101
; CURRENT APPLICATION NUMBER: US/09/498,612
; PRIOR FILING DATE: 2000-02-04
; PRIOR FILING DATE: 1999-02-04
; PRIOR FILING DATE: 1999-02-04
; PRIOR FILING DATE: 1999-02-04
; PRIOR FILING DATE: 1999-03-02
; PRIOR FILING DATE: 1999-03-02
; PRIOR FILING DATE: 1999-03-17
; PRIOR FILING DATE: 1999-03-17
; PRIOR FILING DATE: 1999-04-27
; PRIOR FILING DATE: 1999-04-27
; PRIOR FILING DATE: 2000-02-04
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 6
; LENGTH: 468
; TYPE: PRT
; ORGANISM: Ichthyophthirius multifiliis
US-09-498-612-6

Query Match 100.08; Score 2540; DB 18; Length 468;
 Best Local Similarity 100.08; Pred. No. 3.1e-213;
 Matches 468; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MKNILVILISLFINQISANCPCVGTENTAGQVDDLTGPANCVCQKFNFNNAAFV 60
 DB 1 MKNILVILISLFINQISANCPCVGTENTAGQVDDLTGPANCVCQKFNFNNAAFV 60
 QY 61 PGASTCTPCQKDKAGQNPATANLVTCQNVKCPAGTAIAGGATDYAAIITECVNCRI 120
 DB 61 PGASTCTPCQKDKAGQNPATANLVTCQNVKCPAGTAIAGGATDYAAIITECVNCRI 120
 QY 121 NFYNENAPNAGASTCTACPNVRVGGALTAGNAATIVACQNVACPTGALDDGVTDDYV 180
 DB 121 NFYNENAPNAGASTCTACPNVRVGGALTAGNAATIVACQNVACPTGALDDGVTDDYV 180
 QY 181 RSTFTECVKCLNRYNGNTPFNGKSCQTPCPAIPKPAVQAATLGNDAITTAQCENVA 240
 DB 181 RSTFTECVKCLNRYNGNTPFNGKSCQTPCPAIPKPAVQAATLGNDAITTAQCENVA 240
 QY 241 CPDGTISAAGVNNWVAQNTCTNCAPNFYNNAPNPNPNCSTCLPCPANKDYGAEATAGG 300
 DB 241 CPDGTISAAGVNNWVAQNTCTNCAPNFYNNAPNPNPNCSTCLPCPANKDYGAEATAGG 300
 QY 301 AATLAKOCNCTACPDGTATAGATGATNYVILQTECLNCAANFYDGNFQAGSSRCKACPANK 360
 DB 301 AATLAKOCNCTACPDGTATAGATGATNYVILQTECLNCAANFYDGNFQAGSSRCKACPANK 360
 QY 361 VOGAVATAGGTATLIAQALECPAGTIVLDGTTSTYKQAASECVKCAANFYTTKQTDWVA 420
 DB 361 VOGAVATAGGTATLIAQALECPAGTIVLDGTTSTYKQAASECVKCAANFYTTKQTDWVA 420
 QY 421 GIDTCTSCNKKLTSGAEPANLPESAKNIOQDFANFLSISLLLSIYLL 468
 DB 421 GIDTCTSCNKKLTSGAEPANLPESAKNIOQDFANFLSISLLLSIYLL 468

RESULT 2
 US-09-498-612-5
 ; Sequence 5, Application US/09498612
 ; GENERAL INFORMATION:
 ; APPLICANT: GAERTIG, Jacak
 ; APPLICANT: DICKERSON Jr., Harry W.
 ; APPLICANT: THE UNIVERSITY OF GEORGIA RESEARCH FOUNDATION, INC
 ; TITLE OF INVENTION: RECOMBINANT EXPRESSION OF HETEROLOGOUS NUCLEIC ACIDS IN
 ; FILE REFERENCE: 235.00100101
 ; CURRENT APPLICATION NUMBER: US/09/498,612
 ; CURRENT FILING DATE: 2000-02-04
 ; PRIOR APPLICATION NUMBER: 60/118,634
 ; PRIOR FILING DATE: 1999-02-04
 ; PRIOR APPLICATION NUMBER: 60/122,372
 ; PRIOR FILING DATE: 1999-03-02
 ; PRIOR APPLICATION NUMBER: 60/124,905
 ; PRIOR FILING DATE: 1999-03-17
 ; PRIOR APPLICATION NUMBER: 60/131,121
 ; PRIOR FILING DATE: 1999-04-27
 ; PRIOR APPLICATION NUMBER: PCT/US00/02966
 ; PRIOR FILING DATE: 2000-02-04
 ; NUMBER OF SEQ ID NOS: 14
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO 5
 ; LENGTH: 442
 ; TYPE: PPT
 ; ORGANISM: Ichthyophthirius multifiliis
 US-09-498-612-5

Query Match 36.3%; Score 921; DB 18; Length 442;
 Best Local Similarity 41.8%; Pred. No. 1.3e-71;
 Matches 214; Conservative 45; Mismatches 139; Indels 114; Gaps 19;

QY 1 MKNILVILISLFINQISANCPCVGTENTAGQVDDLTGPANCVCQKFNFNNA 56
 DB 1 MKNILVILISLFINQISANCPCVGTENTAGQVDDLTGPANCVCQKFNFNNGG 56
 QY 57 AAFVPGASTCTPCQKDKAGQNPATANLVTCQNVKCPAGTAIAGGATDYAAIITECV 116
 DB 57 AA-----OGEANGNQPPAAN----- 71
 QY 117 NCRINFENAPNAGASTCTACPNVRVGGALTAGNAATIVACQNVACPTGALDDGVT 176
 DB 72 -----NAARGICVPCQINRVGSVTTNAGDLATLATCSTOCPTGTALDDGVT 117
 QY 177 TDYVRSFTECVKCLNRYNGNTPFNGKSCQTPCPAIPKPAVQAATLGNDAITTAQCENVA 221
 DB 118 DVFDRAAQCCKVCPNFYNGGSPQGEAPVQVFAAGAAAGVAAVTSQCVPCQINKN--N 175
 QY 222 VAQATLGNDAITTAQCNVACPDGTTISAGVNNWVAQNTCTNCAPNFYNN-----N 272
 DB 176 DSPATAGAANLATQCSNQCTGTIVLDGVT--LVFNTSATLVCVCRPNFYNGSPQGE 233
 QY 273 APN-----NSTCLPCPANKDYGAEATAGGATGAAATLAKOCNCTACPDGTATLIAQCA 320
 DB 234 APGVQVFAAGAAAGVAAVTSQCVPCQINKN--DSPATAGAANLATQCSNQCTGTATQD 292
 QY 321 GAT--NYVILQTECLNCAANFYDGNFQAGSSRCKACPANKVQAVATAGGTATLIAQCA 379
 DB 293 GVTLVFSNSTQCSQCIANYFPNG--NFEAGKSQCLKCPVSKTTPAHA--PGNTATQATQCL 350
 QY 380 LECPAAGTIVLDGTTSTYKQAASECVKCAANFYTTKQTDWAGIDTCTSCNKKLTSGAEPAN 439
 DB 351 TPCPAGTIVLDGTTSTYKQAASECVKCAANFYTTKQTDWAGIDTCTSCNKKLTSGAEPAN 439
 QY 440 LPESAKNIOQDFANFLSISLLLSIYLL 468
 DB 411 VYAEATQKVCQASTTFAKFLSISLLLSIYLL 442

RESULT 3
 US-07-763-352A-15
 ; Sequence 15, Application US/07763352A
 ; GENERAL INFORMATION:
 ; APPLICANT: Clark, Theodore G.
 ; APPLICANT: Dickerson, Harry W.
 ; TITLE OF INVENTION: ICH IMMOBILIZATION ANTIGEN AND FISH
 ; TITLE OF INVENTION: VACCINE
 ; NUMBER OF SEQUENCES: 15
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSER: Greenlee and Winner
 ; STREET: 5370 Manhattan Circle, Ste. 201
 ; CITY: Boulder
 ; STATE: Colorado
 ; COUNTRY: USA
 ; ZIP: 80303
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Floppy disk
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: PatentIn Release #1.0, Version #1.25
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/07/763,352A
 ; FILING DATE: 19910920
 ; CLASSIFICATION: 435
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: Ferber, Donna M.
 ; REGISTRATION NUMBER: 33,878
 ; REFERENCE/DOCKET NUMBER: 15-91
 ; TELEPHONE: 303/499-8080
 ; TELEFAX: 303/499-8089
 ; TELEX: 823189
 ; INFORMATION FOR SEQ ID NO: 15:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 414 amino acids

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; TYPE: AMINO ACID
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-07-763-352A-15

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Query Match	33.2%;	Score 843.5;	DB 3;
Best Local Similarity	40.9%;	Pred. NO. 7.3e-65;	Length 414;
Matches 196;	Conservative 41;	Mismatches 131;	Indels 111;
Gaps 18;			

Qy	1	MKNNVILVILISLFTNIKSGANCPVGTEFNTAGQVD----	DLGTPANCVNCKNFYNNNA	56
Db	1	MYNILLVILISLFTNELRVPVCPDGTQ-QAGLTDVGAADLGT----	CVNCRPNFYNGG	56
Qy	57	AAFPVGASTCTPCPKKDGAGQNPBPATANLVTQCNVRCPGATAGGATYAAITCEV	116	
Db	57	AA-----QGEANGNQFPAAN-----	71	
Qy	117	NCRINFYENAPNPNAGASTCTCAPNVRVGGALTAGNAATIVAOQCNVACPTGTALDDGVT	176	
Db	72	-----NAARGICVCPQINRVGSVTNAGDLATLATQCSCTQCTGTALDDGVT	117	
Qy	177	TDVRSFTCEVKRLNFYNGNN--GNTP-----ENPG-----	KSQCTPCPAIKPAN	221
		:		
Db	118	DVFDSSAAQCVKCRPNFYNGSGPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--N	175	
		:		
Qy	222	VAQATLGNDAITTAQCNVACPDGTIIAAGVNNWVAQNT-----CTNCAPNFYNN-----	N	272
		:		
Db	176	DSPATAGAAQANLATQCSNOCPTGTVLDDGVT--LVFNISATLCVKCRPNFYNGSGPQGE	233	
		:		
Qy	273	APN-----FNPG-----NSTCLPCPANKDYGAEATAGGAATLAKOCTACPDGTAIAS	320	
Db	234	APGVQVFAAGAAAAGVAAVTSQCVPCQLNKN--DSPATAGAAQANLATQCSCTQCTGTATD	292	
Qy	321	GAT--NYVILQTECLUNCAANFYFDGNNFQAGSSRCKACPANKVQGVAVATAGGTATLIAQCA	379	
		:		
Db	293	GVTLVFNSSSTQCSOCIANYFFNG--NLEAGKSOCLKCPVSKTTPAHA--PGNTATQATQCL	350	
		:		
Qy	380	LECPAGTVLTDGTSITYKQAASECVKCAANFYTHQTQDWDVAGIDTCTSCHNKILUTSGAEA	438	
		:		
Db	351	TTCPAGTVLDDGTSITNFVASATECTKCSAGTSFKTGTAGTDTCTECTKILTSGATA	409	
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RESULT 4

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US-07-763-352A-3
: sequence 3, Application US/07763352A
: GENERAL INFORMATION:
: APPLICANT: Clark, Theodore G.
: APPLICANT: Dickerson, Harry W.
: TITLE OF INVENTION: ICH IMMOBILIZATION ANTIGEN AND FISH
: TITLE OF INVENTION: VACCINE
: NUMBER OF SEQUENCES: 15
: CORRESPONDENCE ADDRESS:
: ADDRESSEE: Greenlee and Winner
: STREET: 5370 Manhattan Circle, Ste. 201
: CITY: Boulder
: STATE: Colorado
: COUNTRY: USA
: ZIP: 80303
: COMPUTER READABLE FORM:
: MEDIUM TYPE: Floppy disk
: COMPUTER: IBM PC compatible
: OPERATING SYSTEM: PC-DOS/MS-DOS
: SOFTWARE: PatentIn Release #1.0, Version #1.25
: CURRENT APPLICATION DATA:
: APPLICATION NUMBER: US/07/763,352A
: FILING DATE: 19910920
: CLASSIFICATION: 435
: ATTORNEY/AGENT INFORMATION:
: NAME: Ferber, Donna M.
: REGISTRATION NUMBER: 33,878
: REFERENCE/DOCKET NUMBER: 15-91
: TELECOMMUNICATION INFORMATION:

```

```

: TELEPHONE: 303/499-8080
:
: TELEFAX: 303/499-8089
:
: TELEX: 823189
:
: INFORMATION FOR SEQ ID NO: 3:
:
: SEQUENCE CHARACTERISTICS:
:
: LENGTH: 375 amino acids
:
: TYPE: AMINO ACID
:
: TOPOLOGY: linear
:
: MOLECULE TYPE: protein
:
: US-07-763-352A-3

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Query Match	29.5%;	Score 749;	DB 3;	Length 375;
Best Local Similarity	38.4%;	Pred. No. 1.2e-56;		
Matches 172; Conservative	39;	Mismatches 133;	Indels 104;	Gaps 15;

Qy	17	QIKSANCPCVGTETNTAGVD-----DLGTPANCVCOKNFYYNNAAAFVPFGASTCTPCPQK	72
Dd	1	EFRAPCPDGTOTO-AGLTDVGAAADLGT----CNCRPNEYNGGAA-----	42
Qy	73	KDAGAQPNPATANLVTCNVKCPAGTAIAGGATDYAAIIITECVNCRINFNYENAPFNA	132
Dd	43	-QEANGNQPPAAN-----NA	57
Qy	133	GASTCTACPVNRGALTAGNAATIQAQCNVACPTGTALDDGVTTDYVRSETECVKCLRN	192
Dd	58	ARGICVPQINRVGSVTNAGDLATLATOCSTOCTGTALDDGVTDVDRSAACVCKCPN	117
Qy	193	EYNGNNGWTFPECKSOCTPCPAIRPANVAQTGLNDATITAQCNVACPDGTTISAAGVN	252
Dd	118	FYINGG-----SPQGEAPGVQVFAGAAAQAQANLATOCSSNOCTGTVLDDGVT	167
Qy	253	NWVAQNTE---CTNCAPNFYNN-----NAPN---FNPG-----NSTCLPCPANKD	291
Dd	168	--LVENTSATLCVKCRPNFYNGSGSQGEAPGVQVFAGAAAAGAAVTSOCVPCQLKNK	225
Qy	292	YGAETAGATATLAKQNCIACPDGTAIASGAT--NVYLOTECLNCAANFYDPGNPFQGS	350
Dd	226	-DSPATAQAANLATOCSTOCTGTATQDGVTLVSNSSTOCSOCIANYFPNG-NLEAGK	283
Qy	351	SRCACAPANKVOGAVATAGTATLIAQCALECAPGTVLTGDGTTSTYKAQASECVKCAANP	410
Dd	284	SOLCACPKYSKTTPAHA-PGWATATQATCULTTCPAGTVLDGGTSTNEVASATECTKSAGF	342
Qy	411	YTTKTODNWAGIDTCTSCNKLLTSGAEA	438
Dd	343	PASKTGTGTAGTDTCTECTKKLSGATA	370

RESULT 5

```

PCT-US01-06960-2
;
; Sequence 2, Application PC/TUS0106960
; GENERAL INFORMATION:
;
; APPLICANT: The Trustees of Columbia University in the City of New York,
; APPLICANT: et al.
;
; TITLE OF INVENTION: Melanoma Differential Associated Gene-5 (mda-5), Promoter
;
; TITLE OF INVENTION: and Uses Thereof
;
; FILE REFERENCE: 0575/60849-A-PCT
;
; CURRENT APPLICATION NUMBER: PCT/US01/06960
;
; CURRENT FILING DATE: 2001-02-28
;
; NUMBER OF SEQ ID NOS: 3
;
; SOFTWARE: PatentIn version 3.0
;
; SEQ ID NO 2
;
; LENGTH: 3131
;
; TYPE: PRT
;
; ORGANISM: Human
;
PCT-US01-06960-2

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Query Match	9.9%;	Score 251;	DB 1;	Length 3131;
Best Local Similarity	24.7%;	Pred. No. 6.6e-12;		
Matches 119; Conservative	6;	Mismatches 177;	Indels 180;	Gaps 19;

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QY 23 CPVGTETNTAGOVDDLTGPANCVN----- 46
Db 381 CAAGCTTCTAGTTAGACAGCTCTGGATAAGTGCATGGAGGAGAACTGTTGACAATTGA 440
QY 47 ---CCKNFYNNAAAFVPGAST---CTPCPKKDDAGAPNPPATANLVTCNVKCPAGT 99
Db 441 AGACA-----GAAACCGGATGCTGCTGC-----AGAAACAATGGAATGAATCAGGT 489
QY 100 ATAGGATDYAAIITECVNCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVA 159
Db 490 GTAAGA-----GAGCTACTAAAA--AGGATTGTGCAGAAA 522
QY 160 QCNVACPTGTTALDDGVTTDYVRSFTECVKRLNFYNGNNGTTPNPKSQTCPAIPK 219
Db 523 GAAACTGTT-----CTCTGCAATTCGTAATGTT-----CTTC----- 556
QY 220 ANVAQATLGNDAITTAOCNVACPDGTISAAGVNNWVAQNTCT-----NCAPNFYNN 271
Db 557 -GTCAAACAGGAACAATGAACCTTGCCAGAGTTAACAGGCTCTGATTGCTCA----- 609
QY 272 NAFNPNFNGSTCLPCPANKDYGAETAG-GAATLAKQCNACPDGTATIASGATNYVILQT 330
Db 610 -----GAAAGCAATGCAGAGATTGAGAATTTATC--ACAAGTTGATGTT----- 651
QY 331 ECLNCAANFYFDGNNFQAGSSRCKACPAKV-----QGAVATA 368
Db 652 -CCTCAAG---TGGAAGAGCAACTCTTTCAACACAGTTACGCCAAATCTGGAGAAGGA 707
QY 369 GGPATLIAQCALECPAGTVLTGTTSTYKQAAASECVK---AANFYTTKQTDWVAGIDTCT 426
Db 708 GGTCTGGGCA-----TGGAGAA--TAACATCATCAGATCATCTTTTGCAGATTCT 756
QY 427 SC 428
Db 757 TC 758

RESULT 7
US-09-515-363B-2
; Sequence 2, Application US/09515363B
; GENERAL INFORMATION:
; APPLICANT: Fisher, Paul
; TITLE OF INVENTION: Melanoma Differential Associated Gene-5 (mda-5), Promoter and
; TITLE OF INVENTION: Thereof
; FILE REFERENCE: 0575/60849
; CURRENT APPLICATION NUMBER: US/09/515,363B
; CURRENT FILING DATE: 2000-02-29
; NUMBER OF SEQ ID NOS: 3
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 2
; LENGTH: 3131
; TYPE: PRT
; ORGANISM: Human
US-09-515-363B-2

Query Match 9.9%; Score 251; DB 19; Length 3131;
Best Local Similarity 24.7%; Pred. No. 6.6e-12;
Matches 119; Conservative 6; Mismatches 177; Indels 180; Gaps 19;

QY 23 CPVGTETNTAGOVDDLTGPANCVN----- 46
Db 381 CAAGCTTCTAGTTAGACAGCTCTGGATAAGTGCATGGAGGAGAACTGTTGACAATTGA 440
QY 47 ---CCKNFYNNAAAFVPGAST---CTPCPKKDDAGAPNPPATANLVTCNVKCPAGT 99
Db 441 AGACA-----GAAACCGGATGCTGCTGC-----AGAAACAATGGAATGAATCAGGT 489
QY 100 ATAGGATDYAAIITECVNCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVA 159
Db 490 GTAAGA-----GAGCTACTAAAA--AGGATTGTGCAGAAA 522
QY 160 QCNVACPTGTTALDDGVTTDYVRSFTECVKRLNFYNGNNGTTPNPKSQTCPAIPK 219
Db 523 GAAACTGTT-----CTCTGCAATTCGTAATGTT-----CTTC----- 556
QY 220 ANVAQATLGNDAITTAOCNVACPDGTISAAGVNNWVAQNTCT-----NCAPNFYNN 271
Db 557 -GTCAAACAGGAACAATGAACCTTGCCAGAGTTAACAGGCTCTGATTGCTCA----- 609
QY 272 NAFNPNFNGSTCLPCPANKDYGAETAG-GAATLAKQCNACPDGTATIASGATNYVILQT 330
Db 610 -----GAAAGCAATGCAGAGATTGAGAATTTATC--ACAAGTTGATGTT----- 651
QY 331 ECLNCAANFYFDGNNFQAGSSRCKACPAKV-----QGAVATA 368
Db 652 -CCTCAAG---TGGAAGAGCAACTCTTTCAACACAGTTACGCCAAATCTGGAGAAGGA 707
QY 369 GGPATLIAQCALECPAGTVLTGTTSTYKQAAASECVK---AANFYTTKQTDWVAGIDTCT 426
Db 708 GGTCTGGGCA-----TGGAGAA--TAACATCATCAGATCATCTTTTGCAGATTCT 756
QY 427 SC 428
Db 757 TC 758

RESULT 6
US-09-515-363-2
; Sequence 2, Application US/09515363
; GENERAL INFORMATION:
; APPLICANT: Fisher, Paul
; TITLE OF INVENTION: Melanoma Differential Associated Gene-5 (mda-5), Promoter and use
; TITLE OF INVENTION: Thereof
; FILE REFERENCE: 0575/60849
; CURRENT APPLICATION NUMBER: US/09/515,363
; CURRENT FILING DATE: 2000-02-29
; NUMBER OF SEQ ID NOS: 3
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 2
; LENGTH: 3131
; TYPE: PRT
; ORGANISM: Human
US-09-515-363-2

Query Match 9.9%; Score 251; DB 19; Length 3131;
Best Local Similarity 24.7%; Pred. No. 6.6e-12;
Matches 119; Conservative 6; Mismatches 177; Indels 180; Gaps 19;

QY 23 CPVGTETNTAGOVDDLTGPANCVN----- 46
Db 381 CAAGCTTCTAGTTAGACAGCTCTGGATAAGTGCATGGAGGAGAACTGTTGACAATTGA 440
QY 47 ---CCKNFYNNAAAFVPGAST---CTPCPKKDDAGAPNPPATANLVTCNVKCPAGT 99
Db 441 AGACA-----GAAACCGGATGCTGCTGC-----AGAAACAATGGAATGAATCAGGT 489
QY 100 ATAGGATDYAAIITECVNCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVA 159
Db 490 GTAAGA-----GAGCTACTAAAA--AGGATTGTGCAGAAA 522
QY 160 QCNVACPTGTTALDDGVTTDYVRSFTECVKRLNFYNGNNGTTPNPKSQTCPAIPK 219
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Db 523 GAAACTGTT-----CTCTGCAATTCGTAATGTT-----CTTC----- 556
QY 220 ANVAQATLGNDAITTAOCNVACPDGTISAAGVNNWVAQNTCT-----NCAPNFYNN 271
Db 557 -GTCAAACAGGAACAATGAACCTTGCCAGAGTTAACAGGCTCTGATTGCTCA----- 609
QY 272 NAFNPNFNGSTCLPCPANKDYGAETAG-GAATLAKQCNACPDGTATIASGATNYVILQT 330
Db 610 -----GAAAGCAATGCAGAGATTGAGAATTTATC--ACAAGTTGATGTT----- 651
QY 331 ECLNCAANFYFDGNNFQAGSSRCKACPAKV-----QGAVATA 368
Db 652 -CCTCAAG---TGGAAGAGCAACTCTTTCAACACAGTTACGCCAAATCTGGAGAAGGA 707
QY 369 GGPATLIAQCALECPAGTVLTGTTSTYKQAAASECVK---AANFYTTKQTDWVAGIDTCT 426
Db 708 GGTCTGGGCA-----TGGAGAA--TAACATCATCAGATCATCTTTTGCAGATTCT 756
QY 427 SC 428
Db 757 TC 758

RESULT 7
US-09-515-363B-2
; Sequence 2, Application US/09515363B
; GENERAL INFORMATION:
; APPLICANT: Fisher, Paul
; TITLE OF INVENTION: Melanoma Differential Associated Gene-5 (mda-5), Promoter and
; TITLE OF INVENTION: Thereof
; FILE REFERENCE: 0575/60849
; CURRENT APPLICATION NUMBER: US/09/515,363B
; CURRENT FILING DATE: 2000-02-29
; NUMBER OF SEQ ID NOS: 3
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 2
; LENGTH: 3131
; TYPE: PRT
; ORGANISM: Human
US-09-515-363B-2

Query Match 9.9%; Score 251; DB 19; Length 3131;
Best Local Similarity 24.7%; Pred. No. 6.6e-12;
Matches 119; Conservative 6; Mismatches 177; Indels 180; Gaps 19;

QY 23 CPVGTETNTAGOVDDLTGPANCVN----- 46
Db 381 CAAGCTTCTAGTTAGACAGCTCTGGATAAGTGCATGGAGGAGAACTGTTGACAATTGA 440
QY 47 ---CCKNFYNNAAAFVPGAST---CTPCPKKDDAGAPNPPATANLVTCNVKCPAGT 99
Db 441 AGACA-----GAAACCGGATGCTGCTGC-----AGAAACAATGGAATGAATCAGGT 489
QY 100 ATAGGATDYAAIITECVNCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVA 159
Db 490 GTAAGA-----GAGCTACTAAAA--AGGATTGTGCAGAAA 522
QY 160 QCNVACPTGTTALDDGVTTDYVRSFTECVKRLNFYNGNNGTTPNPKSQTCPAIPK 219
Db 523 GAAACTGTT-----CTCTGCAATTCGTAATGTT-----CTTC----- 556
QY 220 ANVAQATLGNDAITTAOCNVACPDGTISAAGVNNWVAQNTCT-----NCAPNFYNN 271
Db 557 -GTCAAACAGGAACAATGAACCTTGCCAGAGTTAACAGGCTCTGATTGCTCA----- 609
QY 272 NAFNPNFNGSTCLPCPANKDYGAETAG-GAATLAKQCNACPDGTATIASGATNYVILQT 330
Db 610 -----GAAAGCAATGCAGAGATTGAGAATTTATC--ACAAGTTGATGTT----- 651
QY 331 ECLNCAANFYFDGNNFQAGSSRCKACPAKV-----QGAVATA 368
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Db 652 -CCTCAAG---TGGAAGACAACTTCTTCAACACAGTTCAGCAAACTCGGAGAAGGA 707
QY 369 GGTATLIAOALCPAGTTLTGTTSTYKQAASECVKC--RANFYTTTQTDWVAGIDTCT 426
Db 708 GGTCTGGGCA-----TGAGAA--TAACATCAGAAATCATCTTTTGCAGATCT 756
QY 427 SC 428
Db 757 TC 758
RESULT 8
US-08-028-021-1
; Sequence 1, Application US/08028021
; GENERAL INFORMATION:
; APPLICANT: THOMPSON, JOHN F.
; TITLE OF INVENTION: PROMOTER AND GENE FOR HUMAN
; TITLE OF INVENTION: CHOLESTEROL
; NUMBER OF SEQUENCES: 7-A HYDROXYLASE
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: PETER C. RICHARDSON
; STREET: 235 EAST 42ND STREET, 42ND STREET
; CITY: NEW YORK
; STATE: NEW YORK
; COUNTRY: U.S.A.
; ZIP: 10017-5755
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/028,021
; FILING DATE: 08-MAR-1993
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: STRASSBURGER, PHILIP C.
; REGISTRATION NUMBER: 34,258
; REFERENCE/DOCKET NUMBER: PC8143
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 573-5731
; TELEFAX: (212) 573-1939
; TELEX: N/A
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 3880 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: DNA (genomic)
US-08-028-021-1
Query Match 9.7%; Score 245.5; DB 4; Length 3880;
Best Local Similarity 25.8%; Pred. No. 2.6e-11;
Matches 115; Conservative 9; Mismatches 198; Indels 123; Gaps 21;
QY 21 ANCPVGTETNAGVDLGTGPANCVNCKNFYNNAAAFVPGASTC-----TPCP 70
Db 2849 ATCATATCTAAA-----GTTGTCCTC-----ATTCCCATAGCTTTCT 2887
QY 71 QKKDAGQPNPAPANLVQC-NVKCPAGTAIAGGATDYAAIITECVNCRINFYENAPN 129
Db 2888 ATACCATGTTTTTATTTTTTTTTCATAACATGATTTTACTCCTCTTCCAT----TGCA 2943
QY 130 FNAGASTCTACPNRVGSGALTAGN-AATIVACNVACPGTALDDGVTTDYVRSFTCEVK 188
Db 2944 ATAGANTCTCCAT--TAGATAGAAATCTGCCTATCTTAT-----TAATG 2987
QY 189 CRLNFYNGNNGNTPFNPGRKSOCTPCPAIK--PANVAQATLGNDAITACNVACPDGPI 246
Db 2988 C-----CTGCACTGGAACTACTTTTGAAGAGTTCTTGGCACGTA 3026

QY 247 SAAGVNNVQAQNTCTNCPNFNNPNFNNPNCSTCLPCPANKDYGAETATGGAATLAK 306
Db 3027 ATAAA-----TACT-CAA-CTAATAATTTTGTGTACACA-----GAAATAAAGTTGG 3072
QY 307 QCNIACPCDGTATASGATNVVILQTECLNCAANFYDGNHNFQAGSSRCACAPANKVQGA 366
Db 3073 AGAACAGATGCCAAT-----TGTACTAG-----TGGTTACTTCTG-----AGTA 3114
QY 367 TAGGTATLIAOALCPAG-----TVLTGDTTSTYKQAASECVKCAANFYTTTQTD 417
Db 3115 AAGGACT--AGCATGTAGTAAATTAATTAATAGATGTTCACTTCCACCAAGATAATGTT 3172
QY 418 VWAG--IDTCT--SCNKKLTSGAEA 438
Db 3173 TTAGTTAGTCTTAACCTTACTTGA 3197
RESULT 9
US-09-791-932-40
; Sequence 40, Application US/09791932
; GENERAL INFORMATION:
; APPLICANT: Vogeli, Gabriel
; APPLICANT: Parodi, Luis A.
; APPLICANT: Hiebsch, Ronald R.
; APPLICANT: Lind, Peter
; APPLICANT: Kaytes, Paul S.
; APPLICANT: Ruff, Valerie
; APPLICANT: Huff, Rita M.
; APPLICANT: Wood, Linda S.
; TITLE OF INVENTION: Novel G Protein-Coupled Receptors Cross-Reference To Related
; FILE REFERENCE: 00325.USI
; CURRENT APPLICATION NUMBER: US/09/791,932
; CURRENT FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/184,305
; PRIOR FILING DATE: 2000-02-23
; PRIOR APPLICATION NUMBER: 60/184,304
; PRIOR FILING DATE: 2000-02-23
; PRIOR APPLICATION NUMBER: 60/184,303
; PRIOR FILING DATE: 2000-02-23
; PRIOR APPLICATION NUMBER: 60/184,397
; PRIOR FILING DATE: 2000-02-23
; PRIOR APPLICATION NUMBER: 60/184,247
; PRIOR FILING DATE: 2000-02-23
; PRIOR APPLICATION NUMBER: 60/188,880
; PRIOR FILING DATE: 2000-03-13
; PRIOR APPLICATION NUMBER: 60/217,369
; PRIOR FILING DATE: 2000-07-11
; PRIOR APPLICATION NUMBER: 60/217,370
; PRIOR FILING DATE: 2000-07-11
; PRIOR APPLICATION NUMBER: 60/218,492
; PRIOR FILING DATE: 2000-07-20
; PRIOR APPLICATION NUMBER: 60/186,810
; PRIOR FILING DATE: 2000-03-03
; PRIOR APPLICATION NUMBER: 60/188,064
; PRIOR FILING DATE: 2000-03-09
; PRIOR APPLICATION NUMBER: 60/186,457
; PRIOR FILING DATE: 2000-03-02
; PRIOR APPLICATION NUMBER: 60/213,861
; PRIOR FILING DATE: 2000-06-23
; PRIOR APPLICATION NUMBER: 60/194,344
; PRIOR FILING DATE: 2000-04-03
; PRIOR APPLICATION NUMBER: 60/218,337
; PRIOR FILING DATE: 2000-07-14
; NUMBER OF SEQ ID NOS: 184
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 40
; LENGTH: 544
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-791-932-40

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Query Match          9.5%; Score 242.5; DB 21; Length 544;
Best Local Similarity 24.5%; Pred. No. 4e-12;
Matches 119; Conservative 11; Mismatches 183; Indels 173; Gaps 22;

QY 23 CPVGTETNTAG-----QVDDLGPANCVNCKNFYNNAAAFVPG-----62
DB 90 CYGGTATTTGGAGCAAAATTTGTACTTTC-----ACAGTTGCTGTGATGSGCA 141
QY 63 -----ASTCTPCPKKDAQAQNPATANILVTCNVKCPAGTAIAGGATDYAAIITEC 115
DB 142 TTTTGTACTCTTC-----TGCTCTCC--ACTTGTG-----CTTC 174
QY 116 VNCRIYNNENAPNENAGASTC-----TACPVNRVGGALTAGNA-----ATIV 158
DB 175 ATC-----TGCAATCGACAGGTACATTTGTGTACTGATCCCTGGCTATGC 221
QY 159 AOCNVA-----CPTGTALDDGVTTDYVRSFTECVKRLNFYNNNGNNTFPNPKSOCT 212
DB 222 TACCAAGTTCCACCGTCTGCTGCGGAATTTGCATCA-----GCGTGTCTCGGATTCT 275
QY 213 PCPAIKPANVAQATLGNDAITTAOCNVACPDGTISAAAGVNNVWVAQNTCTNCAPNFYNN 272
DB 276 GC-----CTCCACGTAC-----ACCGGTGCTGTGTCTACAC-----307
QY 273 APNFPNGNSTCLPCPANKDYGAET--AGGAATLA-----KOCNIAACPD-----GTAIASG 321
DB 308 ---CAGGTGCAATGATGATGGCTGGAGGAATTAAGTCTCTCACTCGGTAGGTG 364
QY 322 -----ATNYILOTECLNCAANFYFDGNFQAGSSRCKACPAKVQAVAT--AGGTA 372
DB 365 GCGTGCAAAATTAATGTAAGTCAAGGCTGGGTGTGTAT-----AGATTTTCTGTGA 414
QY 373 TLIAQCALECPAGTVLTDGTTSTYKQAAAEVCKCAANFYTTKQDVAIGDICTSCNKKL 432
DB 415 TTCTTCACTACCTACCCTGTTGATGATAATTCT-----TTACAGTAAGATTTTCT-TA 466
QY 433 TSGAEA 438
DB 467 TAGCTA 472

RESULT 10
PCT-US97-17746-50
; Sequence 50, Application PC/TUS9717746
; GENERAL INFORMATION:
; APPLICANT: Laten, Howard M.
; TITLE OF INVENTION: PLANT RETROVIRAL POLYNUCLEOTIDES AND
; NUMBER OF SEQUENCES: 58
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Marshall, O'Toole, Gerstein, Murray and Borun
; STREET: 233 South Wacker Drive/6300 Sears Tower
; CITY: Chicago
; STATE: Illinois
; COUNTRY: United States of America
; ZIP: 60606-6402
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US97/17746
; FILING DATE:
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Clough, David W.
; REGISTRATION NUMBER: 36,107
; REFERENCE/DOCKET NUMBER: 27013/33214 US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (312) 474-6300
; TELEFAX: (312) 474-0448
; INFORMATION FOR SEQ ID NO: 50:
```

```
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2826 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
PCT-US97-17746-50

Query Match          9.4%; Score 238.5; DB 1; Length 2826;
Best Local Similarity 26.6%; Pred. No. 7.2e-11;
Matches 119; Conservative 7; Mismatches 229; Indels 93; Gaps 22;

QY 20 SANC-----YGTETINTAGQVDDLGTGPANCV--NCQKNFYNNAAAFVPGASTCTPCPK 73
DB 273 AATCTGTGGTGAATGTCAAGATTGGAAGCAAGTCAAGATGTCCAACCAAGAGCTTC--- 328
QY 74 DAGAQNPPATANILVTCNVKCPAGTAIAGGATDY--AAIITECVNCRINFYNNENAPNF 131
DB 329 -----AACATCAGAC-----CACTCCAGGGTGTGGAACACTCTTCACATGGACITGAT 377
QY 132 AGASTCTA--CPVNRVG---GALTAGNAA-----TIVAOCNVACPTGTALDDGVTTD 178
DB 378 GGGCCCTATGCAAGTTGAAGCCCTTGAAGAAAAGGTATGCCTATGTTGTGGAT-- 435
QY 179 YVRSFTECVKRLNFYNNNGNNTFPNPKSOCTPCPAIKPANVAQATLGNDAITTAOCN 238
DB 436 --GATTTCTCC-----AGATTTACCTGGGTGCAACTT--TATCAGAGAGAAAATCAGACA 484
QY 239 VACPDG-----TISAAGVNNVVAQNT-----ECTNC--APNFYNNAPNFNPNSTCLPCP 287
DB 485 CCTTTGAAGTATTCTCAGAGGTGAGTCTAAGACTTCAAGAGAAAAGACTGTCTCATCA 544
QY 288 ANKDYGAATAGGAATLAKQCNIAACPDGTATAGATNYVILQTECLNCAANFYFDGNFQ 347
DB 545 AGA--GAATCAGGAGTGA-----CCATGGCAGAGAGTTTGAAGAACAGCAAG---TTTACT 594
QY 348 AGSSRCKACPAKVQAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAAAEVCKCA 407
DB 595 GAATTCGTCACTCTGA---AGGCAT---CACTCATGA-----GTTCT-----CTGCA 636
QY 408 ANFYTTKQDVAIGDICTSCNKKLTSG 435
DB 637 GCCATT-----ACACCACACAAAATGG 659

RESULT 11
US-09-254-776-50
; Sequence 50, Application US/09254776
; GENERAL INFORMATION:
; APPLICANT: Laten, Howard M.
; TITLE OF INVENTION: PLANT RETROVIRAL POLYNUCLEOTIDES AND
; NUMBER OF SEQUENCES: 58
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Marshall, O'Toole, Gerstein, Murray and Borun
; STREET: 233 South Wacker Drive/6300 Sears Tower
; CITY: Chicago
; STATE: Illinois
; COUNTRY: United States of America
; ZIP: 60606-6402
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/254,776
; FILING DATE:
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Clough, David W.
; REGISTRATION NUMBER: 36,107
```

REFERENCE/DOCKET NUMBER: 27013/33214 US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 474-6300
TELEFAX: (312) 474-0448
INFORMATION FOR SEQ ID NO: 50:
SEQUENCE CHARACTERISTICS:
LENGTH: 2826 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
US-09-254-776-50

Query Match 9.4%; Score 238.5; DB 16; Length 2826;
Best Local Similarity 26.6%; Pred. No. 7.2e-11;
Matches 119; Conservative 7; Mismatches 229; Indels 93; Gaps 22;

QY 20 SANCP-----VGTETNTAGVDDLTGPANCV--NCQNFYNNAAAFVPGASTCTPCPKK 73
DB 273 AATCTGTGTGAATGTACAGATTGGAAGCAAGTCAAGATGTCACACCAAGCTTC----- 328
QY 74 DAGAQPNNPATANLVTCNVKCPAGTATAGGATDY--AAITECVNCNRYNNAPNFN 131
DB 329 -----AACATCAGAC---CACTTCCAGGGTGTCTGGAAGTACTTACATGGACTTGAT 377
QY 132 AGASTCTA--CPNVNRYG---GALTAGNAA-----TIVACNVACPTGTALDDGVTTD 178
DB 378 GGGGCTATGCAAGTTGAAGCCTTGGAGAAAGGATGATGCCATGTGTGTGGAT-- 435
QY 179 YVRSFTECVKCHLNYNGNNGNTFNPCKSQCTPCPAIKPANVAQATLGNDAITACQN 238
DB 436 --GATTCTCC-----AGATTACCTGGGTCAACTT--TATCAGAGAGAAATCAGACA 484
QY 239 VACPDG-----TISAGYNNVAQNT-----ECTNC-APNFYNNAPNFNPGNSTCLPCP 287
DB 485 CTTTGAAGTATCAAGAGTTGAGTCTAAGACTTCAAGAGAGAAAAAGACTGTGTATCA 544
QY 288 ANKDYGAETAGGAATLAKQCNACPDGTATAGATNIVILQTECLNCAANFYDGNFQ 347
DB 545 AGA--GAATCAGGAGTGA-----CCATGGCAGAGAGTTGAAACACACAAG---TTTACT 594
QY 348 AGSSRCKACPAKPVQAVATAGGTATLIAQCALECPAGTVDGTTSTYKQAAASECVKCA 407
DB 595 GAATCTGCACATCTGA---AGGCAT---CACTCATGA-----GTTCT-----CTGCA 636
QY 408 ANFYTTKQTDWVAGTDTCTSCNKLTSG 435
DB 637 GCATT-----ACACCACACAATATG 659

RESULT 12
PCT-US00-08561-47
; Sequence 47, Application PC/TUS0008561
; GENERAL INFORMATION:
; APPLICANT: HSU, Daniel, K.
; APPLICANT: LIU, Fu-Tong
; APPLICANT: DOWLING, Christopher, A.
; TITLE OF INVENTION: GALECTIN EXPRESSION IS INDUCED IN
; TITLE OF INVENTION: CIRRHOTIC LIVER AND HEPATOCELLULAR CARCINOMA
; FILE REFERENCE: DANHSU.001VPC
; CURRENT APPLICATION NUMBER: PCT/US00/08561
; CURRENT FILING DATE: 2000-03-29
; NUMBER OF SEQ ID NOS: 47
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 47
; LENGTH: 914
; TYPE: PRT
; ORGANISM: human
PCT-US00-08561-47

Query Match 9.4%; Score 237.5; DB 1; Length 914;

Best Local Similarity 25.7%; Pred. No. 2.1e-11;
Matches 123; Conservative 9; Mismatches 208; Indels 139; Gaps 19;
QY 23 CPVGTETNTAGVDDLTGPANCVNCKNFYNNAAAFVPGASTC-----TPCQPKDAG 76
DB 500 CACGCTTCAA-----TGAGAACAAAC-----AGGAGAGTCAATTTGTCATACAA 544
QY 77 AOPNPPATANLVTCNVKCPAGTATAGGATDYAAIITECVNCRINRYNNAPNFNAGAST 136
DB 545 AGCTGGATA-----ATAACTGGGGAAGGAAGAACAGACAGTC-----GGTT 585
QY 137 CTACPNRVGGAL-TAGNA-----ATIVACNVACPTGTALDDGVTTDYVRSFTECVKRL 191
DB 586 TTCCTCATTTGAAAGTGGGAAACCATTTCAAAATACAACTACTGGTTGAACCTGACCA----- 641
QY 192 NFYNGNNGNTFNPCKSQCTPCPAIKPANVAQATLGNDAITACNVACPDGTISAGV 251
DB 642 -----CTTCAAGTTGCGATGAATGAT---GCTCCTGTGTGCGAT 679
QY 252 -----NNWVAQNTNCTNCAPNFYNNAPNFNPGNSTCLPCPANKDYGAET--- 297
DB 680 ACAATCATCGGTTTAAATAACT-CAA-----TGAATCAGCAAACTGGGAATTC 728
QY 298 ---AGGAATLAKQCNACPDG-----TATAGATNTVILQTECLNCAANFYDGNFQ 347
DB 729 TGTGACATAGACCTCAGCAGTGTTCATATACCATGATATAATCTGAAA-----GGGC 783
QY 348 AGSSRCKACPAKPVQAVATAGGTATLIAQCALECPAG-----TVLTDGT----- 392
DB 784 AGATTAAAAAAGAAAGAACTTAAACCTTACATGTGTAAAGGTTTATGTTTCACTG 843
QY 393 -----TSTYKQAAASECVKCA--ANFYTTKQTDWVAG-IDTCTSCNKLTSAGEA 438
DB 844 TGATGAAAAATTTTACATTCATCAATCAATCCCTCTCTGTGAAGTCACTACTTAATAATA 902

RESULT 13
US-09-436-063-5
; Sequence 5, Application US/09436063A
; GENERAL INFORMATION:
; APPLICANT: Bamber, Bruce A.
; APPLICANT: Jorgensen, Erik M.
; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and
; TITLE OF INVENTION: Methods Related thereto
; FILE REFERENCE:
; CURRENT APPLICATION NUMBER: US/09/436,063A
; CURRENT FILING DATE: 1999-11-08
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 5
; LENGTH: 1917
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-436-063-5

Query Match 9.2%; Score 233; DB 18; Length 1917;
Best Local Similarity 25.4%; Pred. No. 1.3e-10;
Matches 115; Conservative 10; Mismatches 238; Indels 90; Gaps 18;
QY 23 CPVGTETNTAGVDDLTGPANCVNCKNFYNNAAAFVPGA-----STCTPCPQ 71
DB 622 CTCACCTACCGTCGGAGTAGACCTGGATAGACTGTGGAACCCGACACGCTTCTCC 681
QY 72 KKDAGAPNPATANLVTCNVKCPAGTATAGGATDYAAIITECVNCRINRYNNAPNFN 131
DB 682 AAATGAAA-AGAAATCATCTTCTCCACTT---GGCAACCCACACATAAC-----T 725
QY 132 AGASTCTACPNRVGGALTAGNAATIVACQN-----VACPTGALDDGVTTDYVRSFTEC 186
DB 726 CGTTCCTTCGTATCGAG--GGTATGGAACGGTTTATAGTAGTCAAGATTAAACAGTCAC 783
QY 187 VKRLNFYNGNNGNTFNPCKSQCTPCPAIKPANVAQATLGNDAITIT-AQC-----NVA 240

Wed Dec 26 12:01:34 2001

us-09-497-967-7.rapm

Db 784 TGCAA-----CGTGTCCAATGGACCTGAAGCTGTTCCTCCAAATGGACTCTCAACACTGTATAA 838
QY 241 CPDGTISAAGVNNWVAQNTCTNCAPNFYNNAPNPNPNCNSTCLPCPANKDYGAETAGG 300
Db 839 CTGGAATTTGAAAGCTACGGCTACA-----GTAICCTCGA-----CATTATG 880
QY 301 AATLAKQCNIACPDGTAIASG---ATNYVILQTECLNCAANFYFDGNFQAGSSRCKACP 357
Db 881 TAGCTGTCCGACGAGAAGAGTCCGCG---TCCACCGAGTCTTATG--AGTTGCCGCA 933
QY 358 ANKVOGAVAT-----AGGT-ATLIAQCALEC---PAGTVLTDGTTSTYKQAASEC 403
Db 934 GTTTGACTTCACTCATCAAGTCTCAATCATAGCGAAAAGCTTAGTTCAGGAGAATA 993
QY 404 VKCAANFYTTKQTDWVAGIDTCTSCNKKLTSGA 436
Db 994 TTCCCGCCT---TTGCTGGTCTCTCTATTCAA 1023

RESULT 14
US-09-436-063C-5
; Sequence 5, Application US/09436063C
; GENERAL INFORMATION:
; APPLICANT: Bamber, Bruce
; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and
; FILE REFERENCE: P-1095corrected
; CURRENT APPLICATION NUMBER: US/09/436,063C
; PRIOR FILING DATE: 1999-11-08
; PRIOR FILING DATE: 1998-11-09
; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 5
; LENGTH: 1917
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-436-063C-5

Query Match 9.2%; Score 233; DB 18; Length 1917;
Best Local Similarity 25.4%; Pred. No. 1.3e-10;
Matches 115; Conservative 10; Mismatches 238; Indels 90; Gaps 18;

QY 23 CPVGTETNTAGQVDDLGTPANCVCQKNFYNNAAAFVPGA-----STCTPCPQ 71
Db 622 CTCACCTTACCGTCGGAGTAGACTACCTGGATAGACTGTGGAACCCGACACGTTCTTCCC 681
QY 72 KKDAGAPNPATANLVTCQNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFEN 131
Db 682 AAATGAAA-AGAAATCATCTTCCACTT---GGCAACCACACATAAC-----T 725
QY 132 AGASTCTACPVNRVGGALTAGNAATIVAQCN-----VACPTGTALDDGVTTDYVRSFTEC 186
Db 726 CGTTCCTTCGTATCGAG--GGTGATGGAACGGTTTATATACTAGTCAAAGATTAAACAGTCAC 783
QY 187 VKRLNFYNGNNGNTPNPGKSQCTPCPAIKPANVAQATLGNDAITIT-AQC-----NVA 240
Db 784 TGCAA-----CGTGTCCAATGGACCTGAAGCTGTTCCTCAATGGACTCTCAACACTGTATAA 838
QY 241 CPDGTISAAGVNNWVAQNTCTNCAPNFYNNAPNPNPNCNSTCLPCPANKDYGAETAGG 300
Db 839 CTGGAATTTGAAAGCTACGGCTACA-----GTATCCTCGA-----CATTATG 880
QY 301 AATLAKQCNIACPDGTAIASG---ATNYVILQTECLNCAANFYFDGNFQAGSSRCKACP 357
Db 881 TAGCTGTCCGACGAGAAGAGTCCGCG---TCCACCGAGTCTTATG--AGTTGCCGCA 933
QY 358 ANKVOGAVAT-----AGGT-ATLIAQCALEC---PAGTVLTDGTTSTYKQAASEC 403
Db 934 GTTTGACTTCACTCATCAAGTCTCAATCATAGCGAAAAGCTTAGTTCAGGAGAATA 993

Search completed: December 26, 2001, 10:38:06
Job time: 187 sec

QY 404 VKCAANFYTTKQTDWVAGIDTCTSCNKKLTSGA 436
Db 994 TTCCCGCCT---TTGCTGGTCTCTCTATTCAA 1023
RESULT 15
US-09-627-650-5
; Sequence 5, Application US/09627650
; GENERAL INFORMATION:
; APPLICANT: Bamber, Bruce
; TITLE OF INVENTION: Nematode Neuromuscular Junction GABA Receptors and
; FILE REFERENCE: P-1095corrected
; CURRENT APPLICATION NUMBER: US/09/627,650
; PRIOR FILING DATE: 2000-07-28
; PRIOR FILING DATE: 1998-11-09
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 5
; LENGTH: 1917
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-09-627-650-5

Query Match 9.2%; Score 233; DB 20; Length 1917;
Best Local Similarity 25.4%; Pred. No. 1.3e-10;
Matches 115; Conservative 10; Mismatches 238; Indels 90; Gaps 18;
QY 23 CPVGTETNTAGQVDDLGTPANCVCQKNFYNNAAAFVPGA-----STCTPCPQ 71
Db 622 CTCACCTTACCGTCGGAGTAGACTACCTGGATAGACTGTGGAACCCGACACGTTCTTCCC 681
QY 72 KKDAGAPNPATANLVTCQNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFEN 131
Db 682 AAATGAAA-AGAAATCATCTTCCACTT---GGCAACCACACATAAC-----T 725
QY 132 AGASTCTACPVNRVGGALTAGNAATIVAQCN-----VACPTGTALDDGVTTDYVRSFTEC 186
Db 726 CGTTCCTTCGTATCGAG--GGTGATGGAACGGTTTATATACTAGTCAAAGATTAAACAGTCAC 783
QY 187 VKRLNFYNGNNGNTPNPGKSQCTPCPAIKPANVAQATLGNDAITIT-AQC-----NVA 240
Db 784 TGCAA-----CGTGTCCAATGGACCTGAAGCTGTTCCTCAATGGACTCTCAACACTGTATAA 838
QY 241 CPDGTISAAGVNNWVAQNTCTNCAPNFYNNAPNPNPNCNSTCLPCPANKDYGAETAGG 300
Db 839 CTGGAATTTGAAAGCTACGGCTACA-----GTATCCTCGA-----CATTATG 880
QY 301 AATLAKQCNIACPDGTAIASG---ATNYVILQTECLNCAANFYFDGNFQAGSSRCKACP 357
Db 881 TAGCTGTCCGACGAGAAGAGTCCGCG---TCCACCGAGTCTTATG--AGTTGCCGCA 933
QY 358 ANKVOGAVAT-----AGGT-ATLIAQCALEC---PAGTVLTDGTTSTYKQAASEC 403
Db 934 GTTTGACTTCACTCATCAAGTCTCAATCATAGCGAAAAGCTTAGTTCAGGAGAATA 993
QY 404 VKCAANFYTTKQTDWVAGIDTCTSCNKKLTSGA 436
Db 994 TTCCCGCCT---TTGCTGGTCTCTCTATTCAA 1023